CLAIMS :

- (1.) A cloned DNA which contains a DNA which is hybridizable with the genomic RMA of the LAV viruses on a fragment of said hybridizable DMA.
- 2. The DNA of claim t which is a recombinent of said hybridizable DNA or DNA fragment hybridizable with the general RNA of the LAV virus.
 - 3. The DNA of claim 1 or 2 wherein said hybridizablo DNA or DNA fragment is a cDMA.
- 14. The DHA of claims 1 to 3 which contains the following restriction sites in the following order (from 10 the 3' and to the 5' end) :

Hind III, Sac I, Pgl II /LAV 751.

5. The DMA of claim 4 which contains the following restriction sites in the following order :

Hind III, Sac I, 893/II, 893 II, Kpn I (LAV 82). 15 6. The DNA of claim Wilet contains the following restriction sites in the policwing order :

Hind III, Sec I. Adl II. Bol II, Kon I, XHo I. Bom HI, Hind II/, Bgl II (LAV 13).

7. The DNA of claim 6 which has a size of about 2.5 kb.

- The DNA/of any of claims 1 to 7 which contains a region corresponding to the R and U3 regions of the LTR as well as to /the 3' end of the coding region of the rotroviral ONA.
- 9. The DNA of claim 1 which has a size from about 9.1 to 9.2 kb./
- 10. The DNA of claim 9 which centains the following series of restriction sites:

30	Hind III	0
-	Sa/c I	50
	Bam KI	4 G D
	#ind III	520
	/Bam HI	600
35	/ Pst I	000

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1 100 Hind III 1 500 Dgl II 3 500 Kpn I 3 900 Kpn I 4 10.0 Eco RI 5 3/00 Eco RI 5 5/500 Sal I Kpn I 6 100 6 500 Bgl II 7 600 Bgl III 7 850 Kind III 10 8 150 · Bam HI 8 600 Xho I 8 700 Kpn I 8 750 Bg1 I Bgl I 9 150 15 9 200 Sac I 9 250 Hind III

11. The DNA of claim 10 which contains an additional Hind/III approximately at the 5 550 coordinate.

12. A DNA fragment according to claim (which comprises a sequence extending from approximately Kon I (6100) to approximately Bam HI (8150) of the sequence defined in claim 11.

prises a sequence extending from approximately Kpn 1 (3500) to approximately Bgl II (6500) of the sequence defined in claim 11.

prises a sequence extending from approximately Pst (800) to approximately Kpn I (3500) of the sequence defined in claim 11.

15. A DNA fragment of claim 1 which codes for the envelopes proteins.

15. A DNA fragment of claim 1 which codes for the retroviral polymerase.

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Droteins. ONA fragment which codes for the cor

- 18. A probe for the <u>in vitro</u> detection of LAV which consists of a DNA according to any of claims 1 to 17.
- for the transformation of procaryotic or eucaryotic cells which contains an insert consisting of a DNA fragment hydridizable with the retroviral genome of LAV viruses as defined in any of claims 1 to 17.
 - 20. The vector of claim 18 which contains the DNA fragment of claim 15.
 - 21. A microorganism, eucaryotic or procaryotic cell which is transformed by a vector according to claim 19 or 20 and which expresses the polypeptide encoded by the corresponding DNA fragment.

(22) The purified RNAs of LAV viruses which have salzes from 9.1 to 9.2 kb.

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